FIGURE 1A: V_L DOMAIN

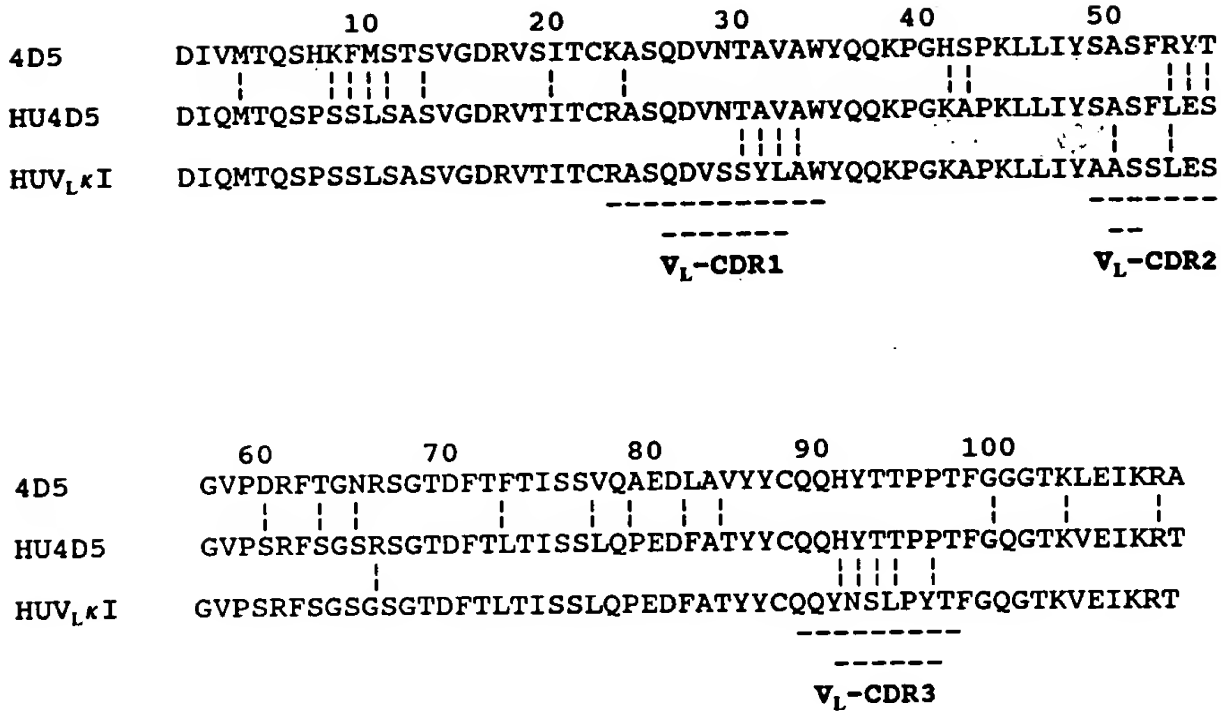


FIGURE 1B: V_H DOMAIN

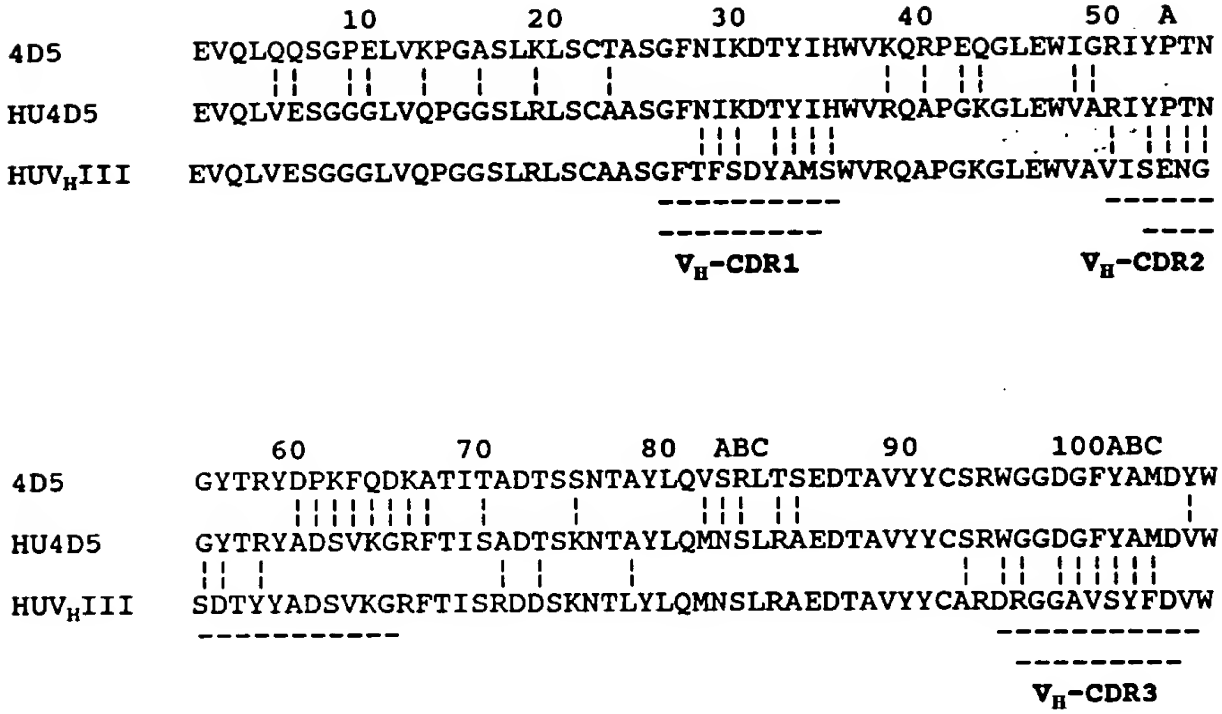
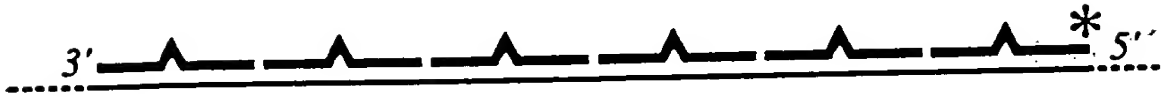
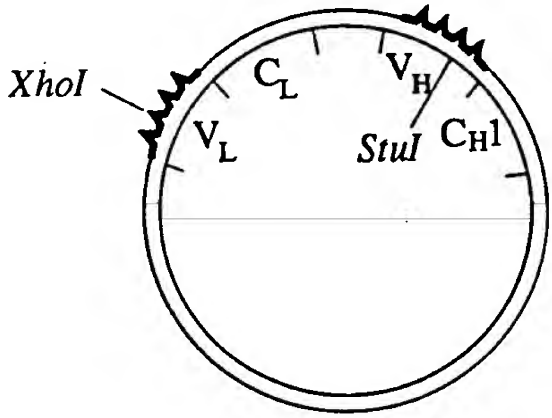


FIGURE 2

Anneal huV_L or huV_H oligomers to pAK1 template



1. Ligate
2. Isolate assembled oligomers
3. Anneal to pAK1 template (*XhoI*$^-$, *StuI*$^+$)
4. Extend and ligate



1. Transform *E. coli*
2. Isolate phagemid pool
3. Enrich for huV_L and huV_H(*Xho I*$^+$, *StuI*$^-$)
4. Sequence verify

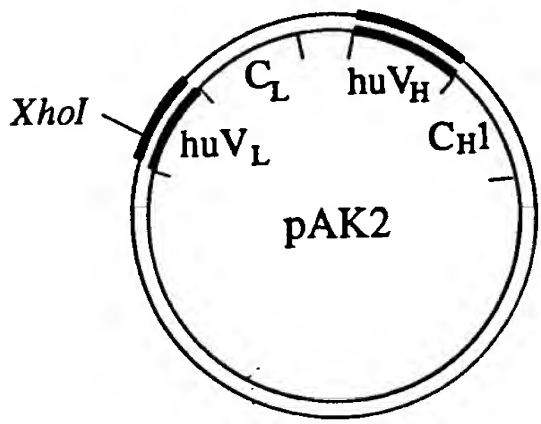


FIGURE 3

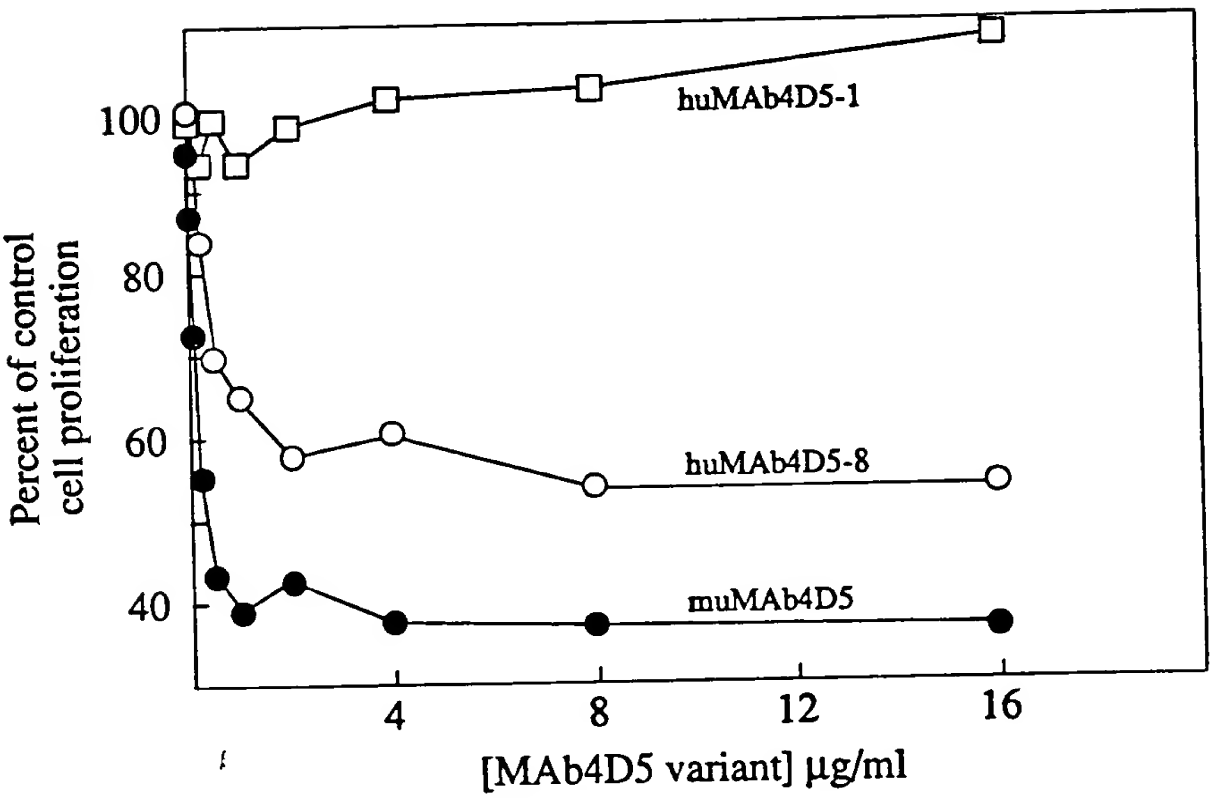

Percent of control cell proliferation
100
80
60
40
huMAb4D5-1
huMAb4D5-8
muMAb4D5
4
8
12
16
[MAb4D5 variant] µg/ml

FIGURE 4